

## Family *Pluralincolimonadaceae*

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### Etymology

[Plu.ral.in.co.li.mo.na.da'ce.ae] N.L. fem. n. *Pluralincolimonas*, referring to the type genus *Pluralincolimonas*; -*aceae*, ending to denote a family; N.L. fem. pl. n. *Pluralincolimonadaceae*, the *Pluralincolimonas* family

### Nomenclatural type

Genus *Pluralincolimonas*

### Description

This family belongs to the order *Pluralincolimonadales*, class *Gorgyria*, phylum *Omnitrophota*. This family is currently a monotypic family, and has the same description as the nomenclatural type of the family, the genus *Pluralincolimonas*.

### Classification

*Bacteria* » *Omnitrophota* » “*Gorgyria*” » *Pluralincolimonadales* » *Pluralincolimonadaceae*

### References

Effective publication: Seymour et al., 2023 [1]

### Registry URL

<https://seqco.de/i:23730>

## References

1. Seymour et al. (2023). Hyperactive nanobacteria with host-dependent traits pervade Omnitrophota. *Nature Microbiology*. [DOI:10.1038/s41564-022-01319-1](https://doi.org/10.1038/s41564-022-01319-1)